

# **Pharmacogenomics: Science and Translation**

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# Pharmacogenomics

- **Introduction**
- **Pharmacogenomic science**
- **Pharmacogenomic translation**
- **Conclusions**

# **Pharmacogenetics-Pharmacogenomics**

**The study of the role of  
inheritance in individual  
variation in drug response  
phenotypes.**

# **Pharmacogenetics-Pharmacogenomics**

## **Clinical Goals**

- Avoid adverse drug reactions
- Maximize drug efficacy
- Select responsive patients

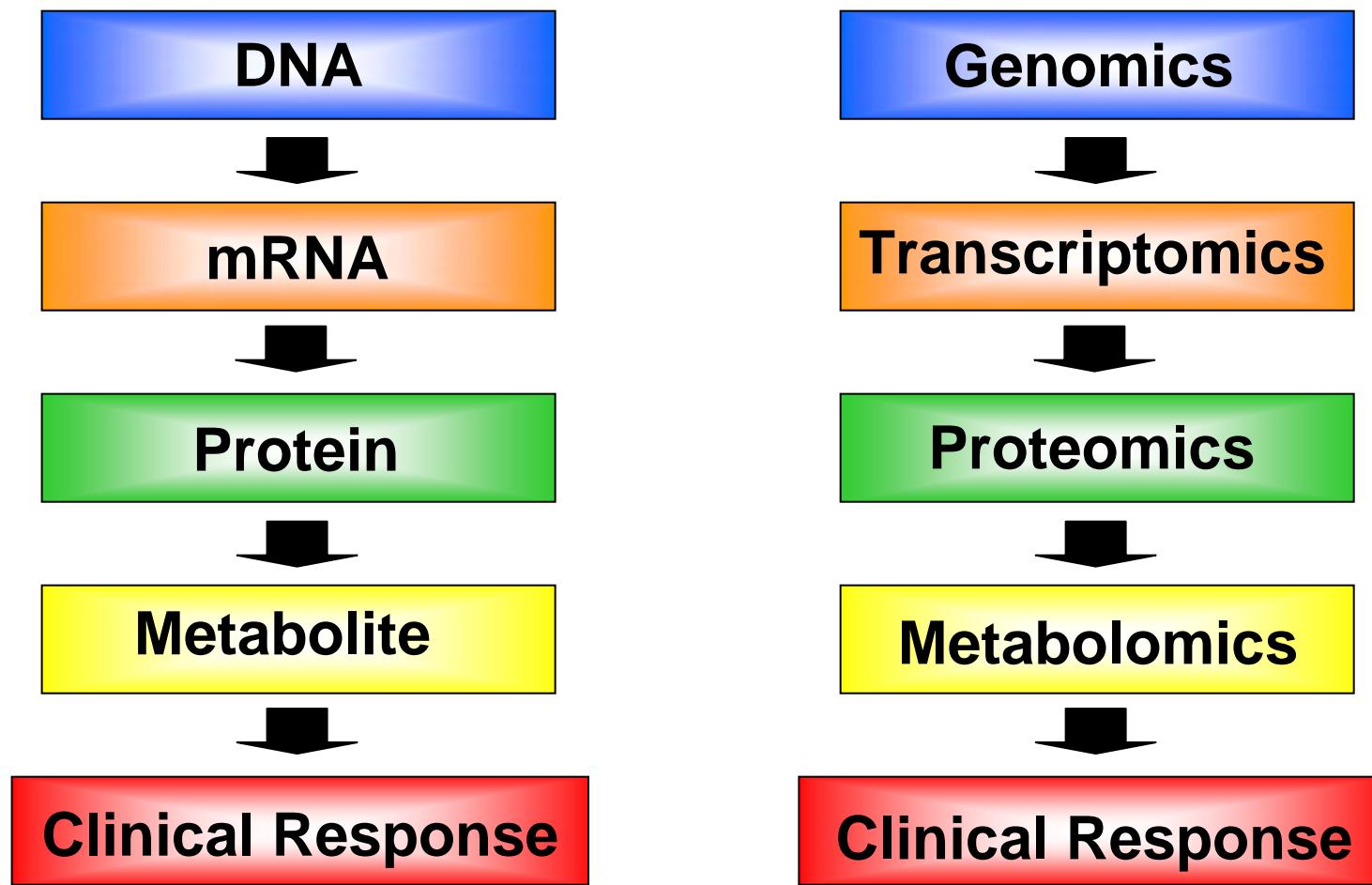
# **Pharmacogenetics-Pharmacogenomics**

## **Scientific Goals**

- Link variation in genotype to variation in phenotype
- Determine mechanisms responsible for that link
- Translate the link into enhanced understanding, treatment and prevention of disease

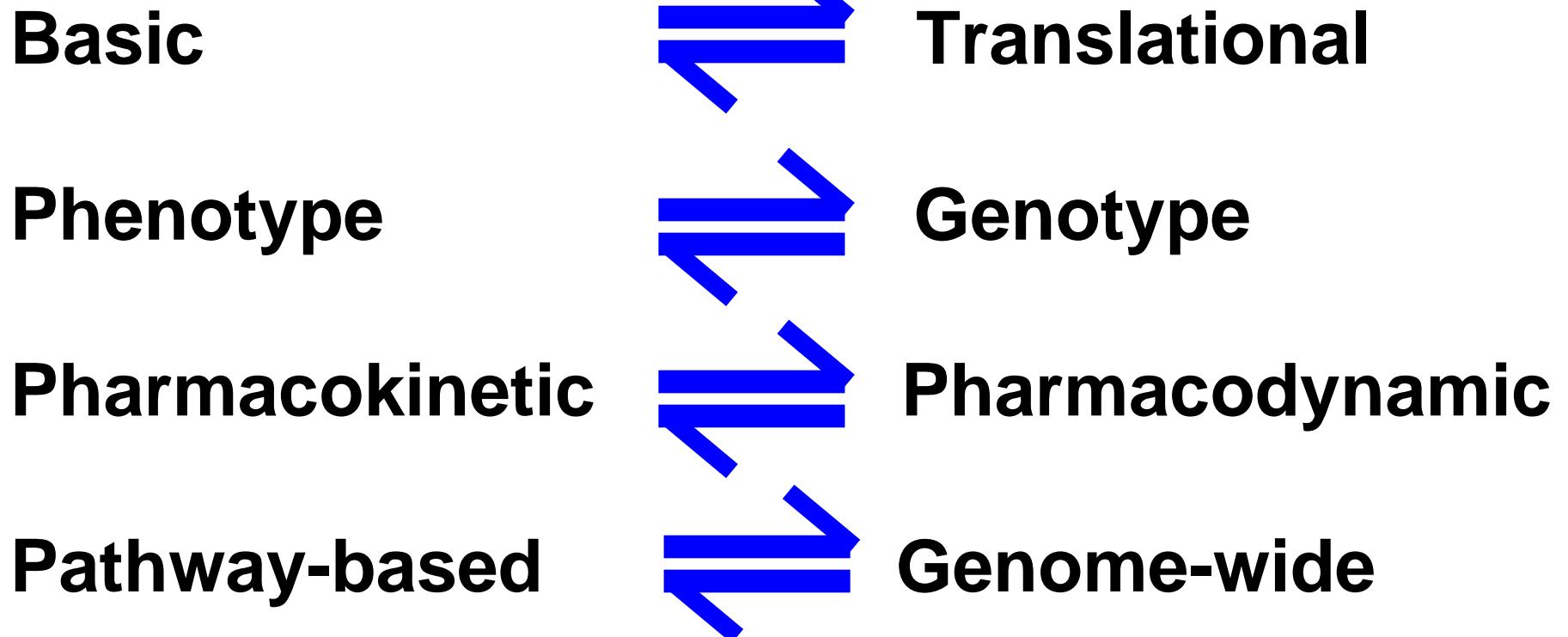
# Pharmacogenetics-Pharmacogenomics

## Drug Response Predictors



# **Pharmacogenetics-Pharmacogenomics**

## **Complementary Approaches**



### NIH Funding Institutes

**NIGMS**  
**NHLBI**  
**NIDA**  
**NCI**  
**NIEHS**  
**NIMH**  
**NHGRI**  
**NLM**  
**ORWH**



● Primary Investigator Site  
● Co-Investigator Site

# Pharmacogenomics

- Introduction
- Pharmacogenomic science
- Pharmacogenomic translation
- Conclusions

# Pharmacogenetics-Pharmacogenomics

## FDA Hearings

## Pharmacogenetics and Drug Labeling

- **Thiopurines – *TPMT***
- **Irinotecan – *UGT1A1***
- **Warfarin – *CYP2C9* and *VKORC1***
- **Tamoxifen – *CYP2D6***

2003 FDA  
“Draft Pharmacogenomic  
Guidance”

Valid Biomarkers

- Thiopurine S-methyltransferase (TPMT)
- Cytochrome P450 2D6 (CYP2D6)

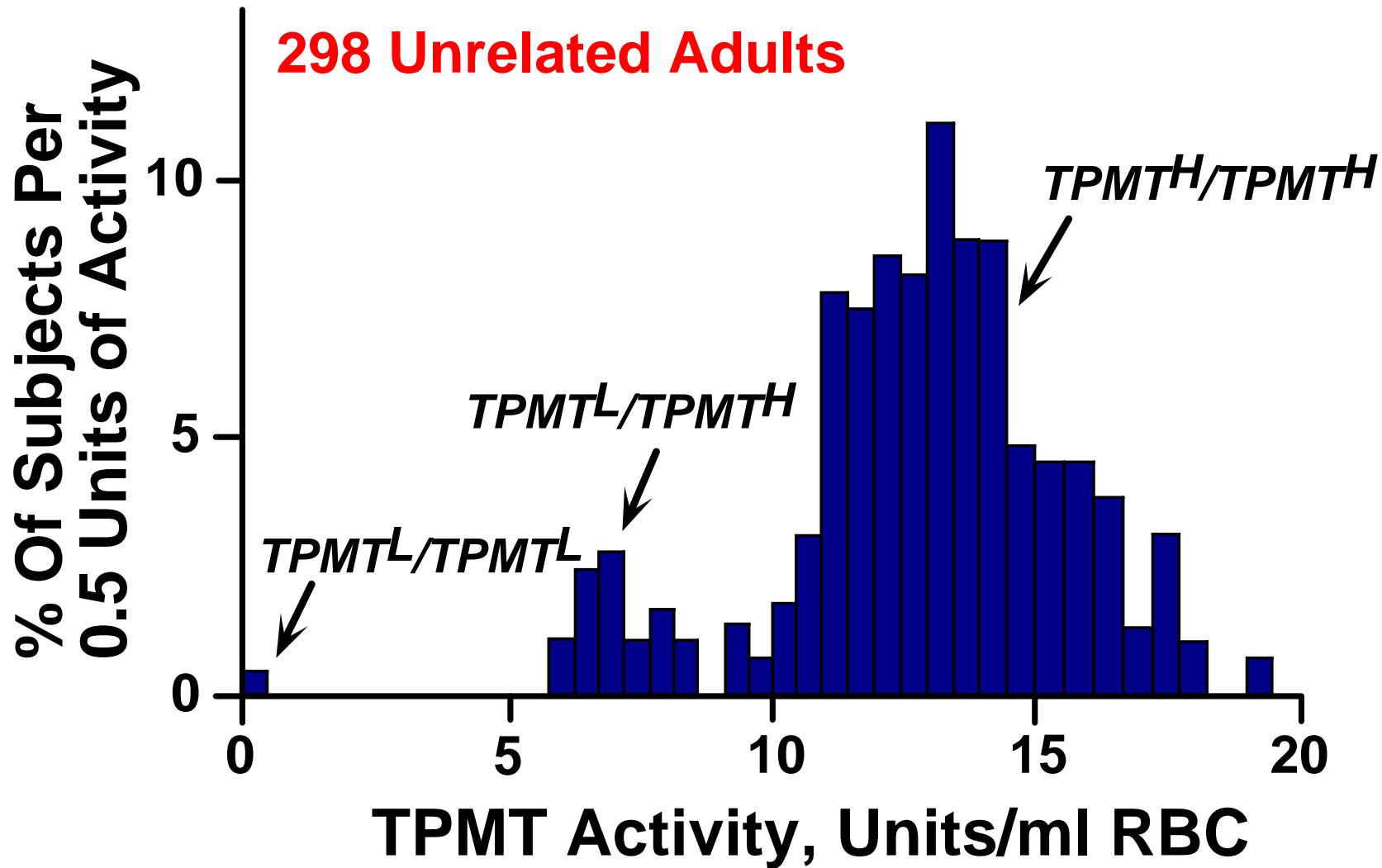
# **TPMT**

# **Pharmacogenomics**

## **Basic Pharmacogenomics**



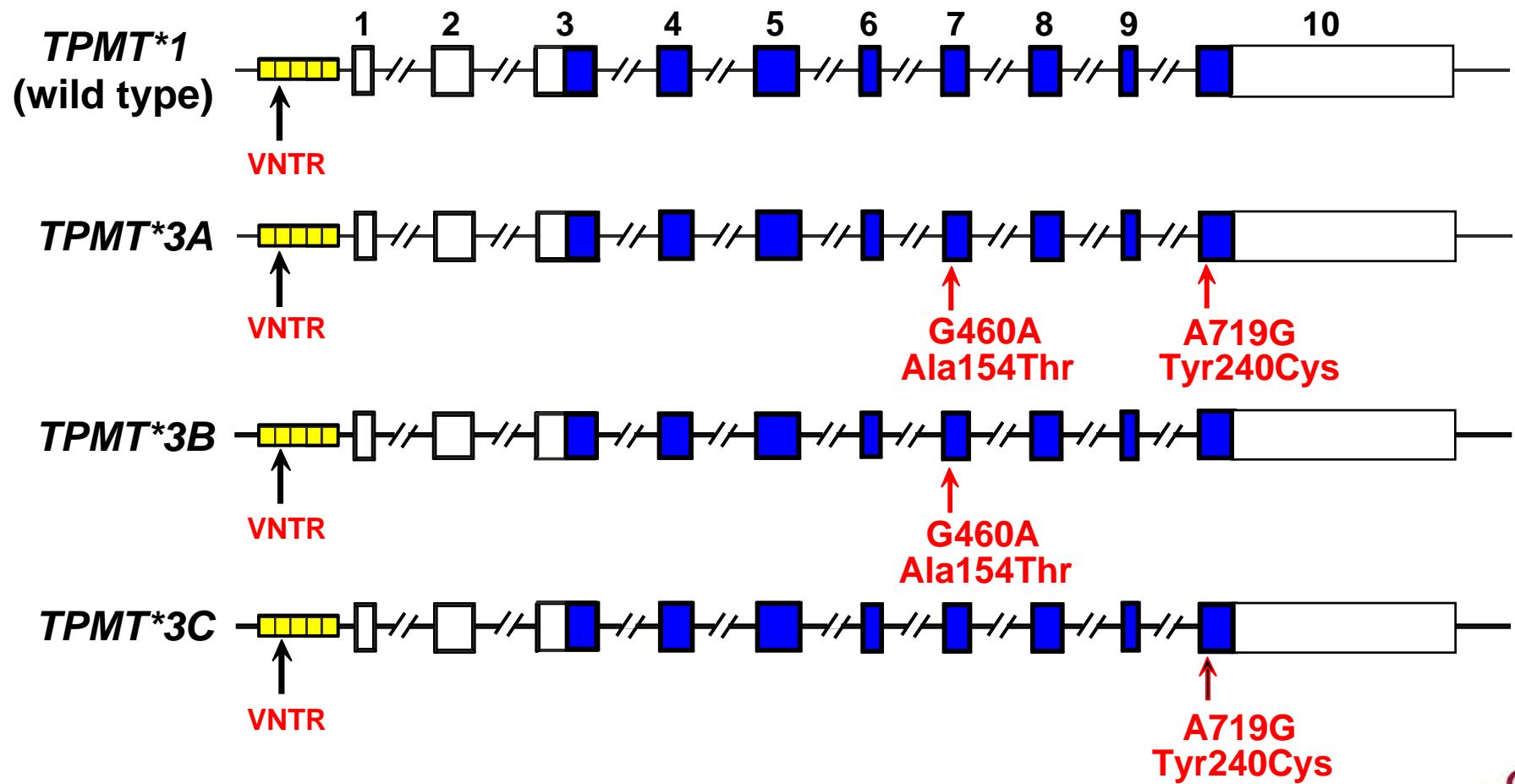
# Human RBC TPMT



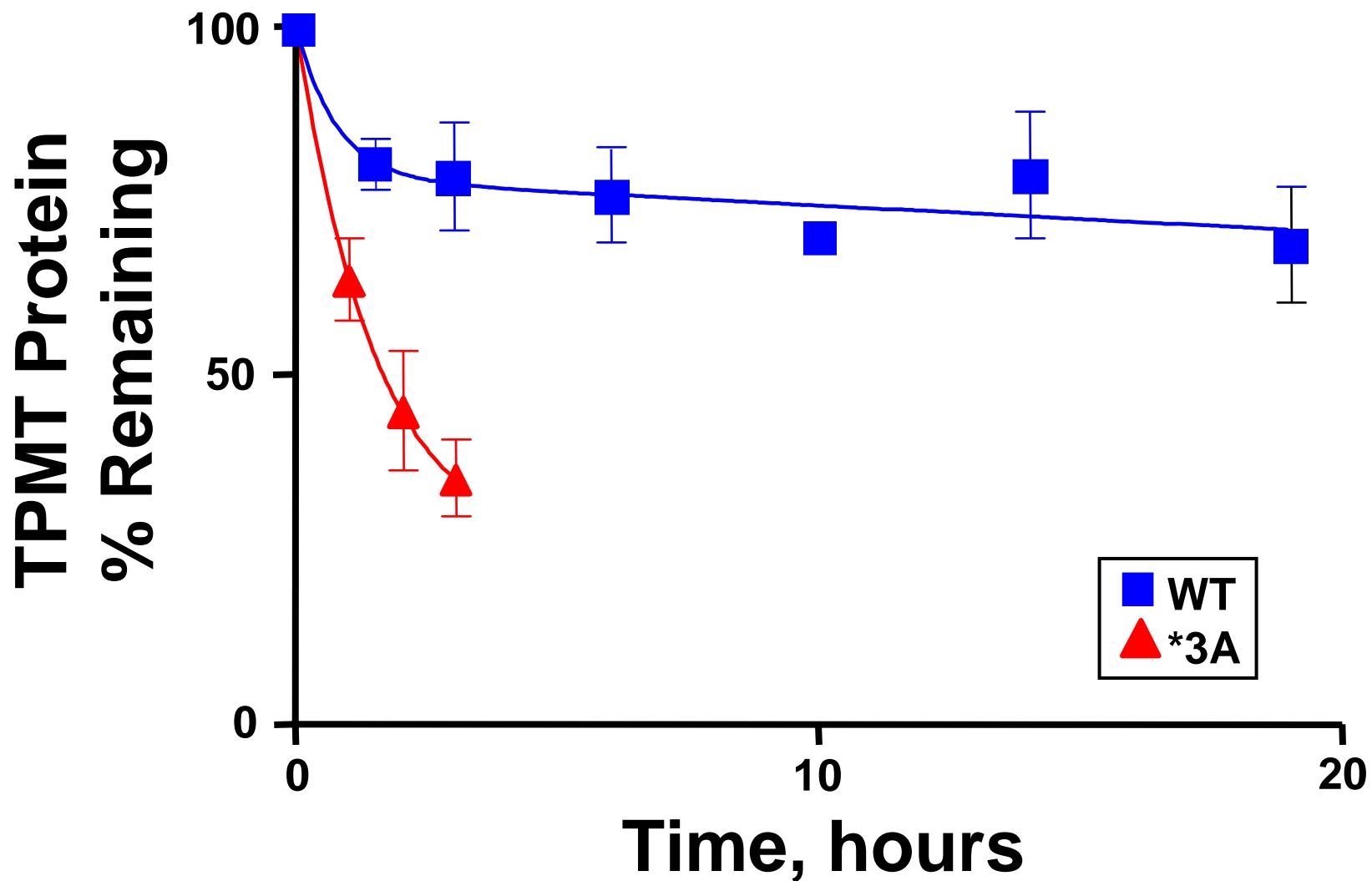
# **TPMT Genetic Polymorphism Clinical Consequences**

- **Low TPMT**
  - Increased thiopurine toxicity
  - Increased risk for secondary neoplasm
- **High TPMT**
  - Decreased therapeutic effect

# Selected Human TPMT Alleles

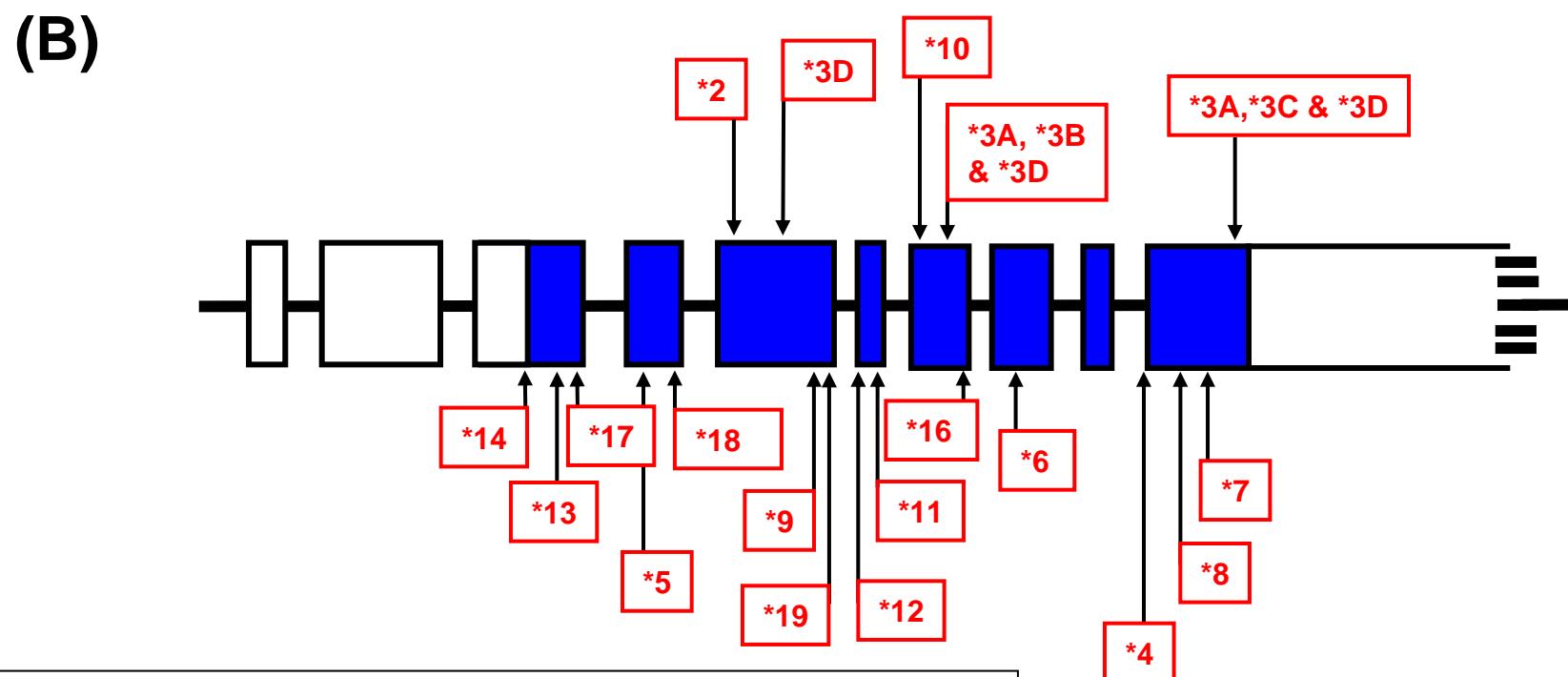
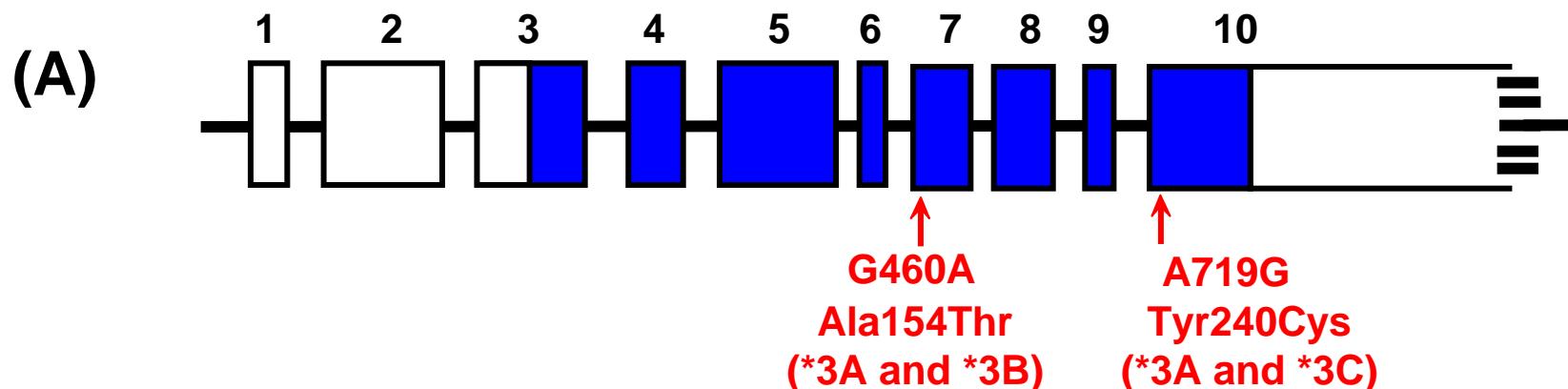


# Human TPMT Protein RRL Degradation



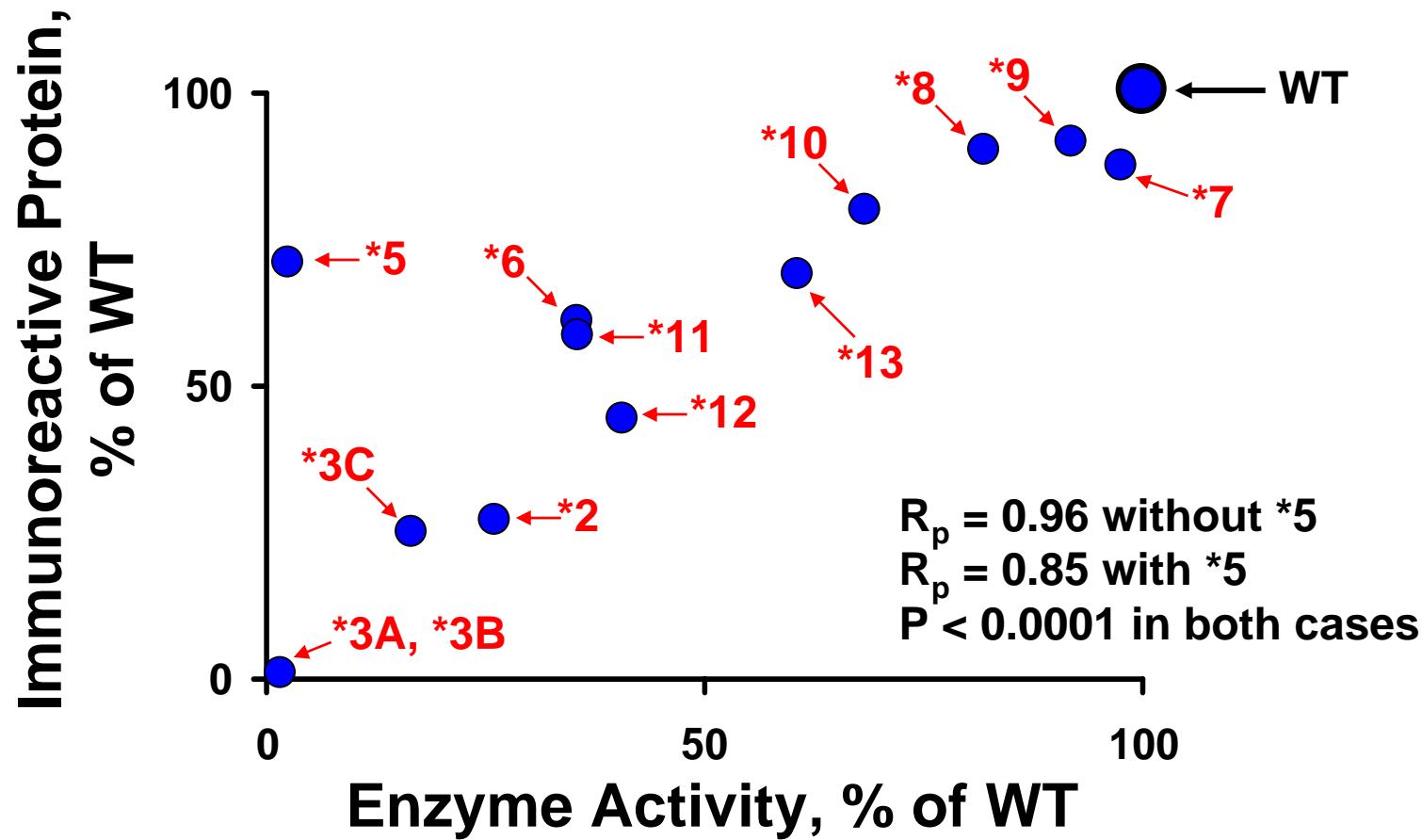
Wang et al., Pharmacogenetics 13:555-64, 2003

# *TPMT* Gene Structure and Polymorphism/Mutation Locations



Salavaggione et al., Pharmacogenet. Genomics 15:801-15, 2005

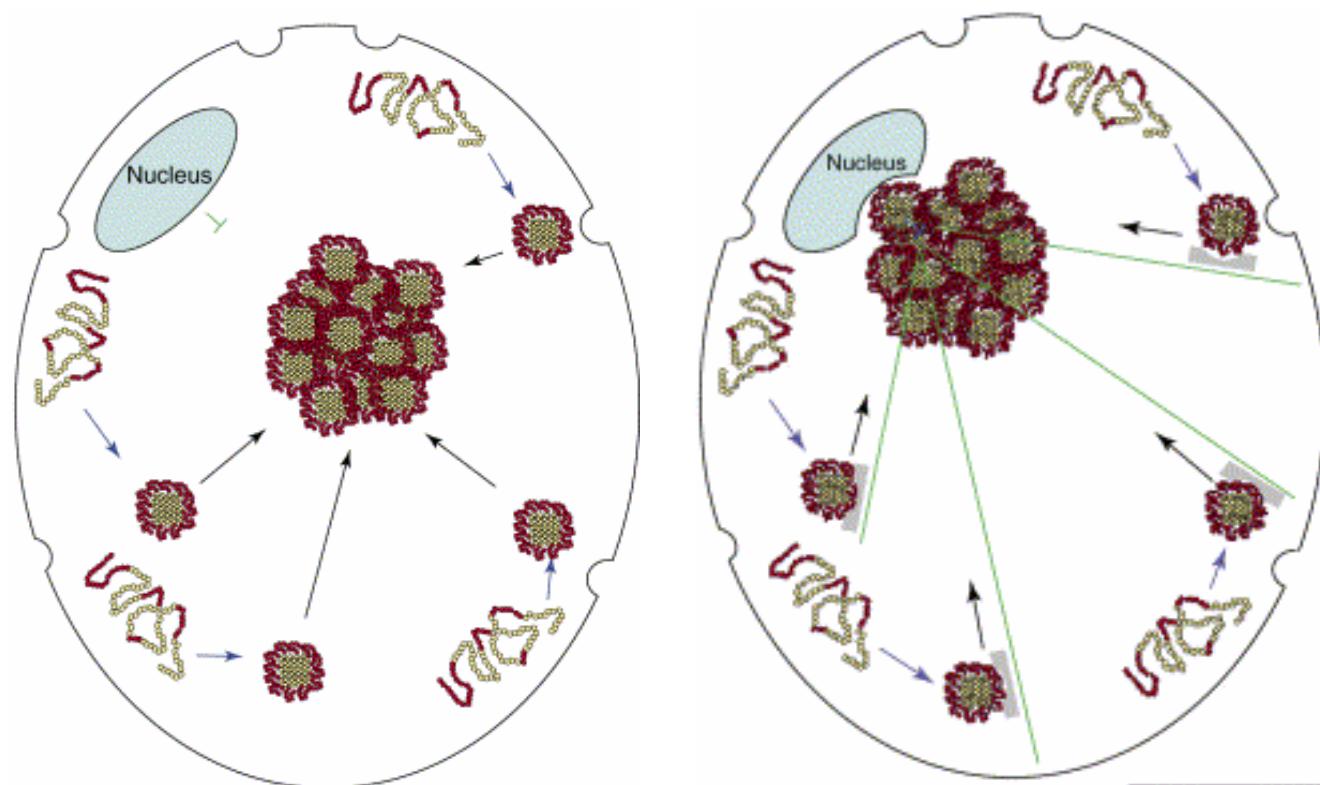
# Recombinant Human TPMT Allozyme Activity and Protein Levels



# Hypothesis

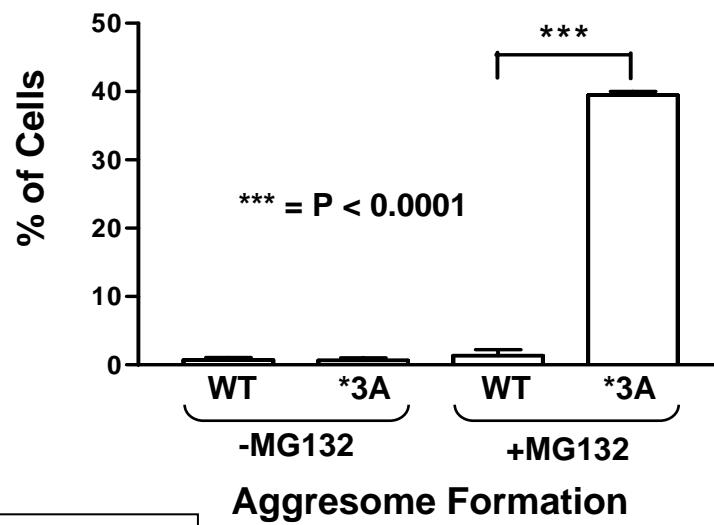
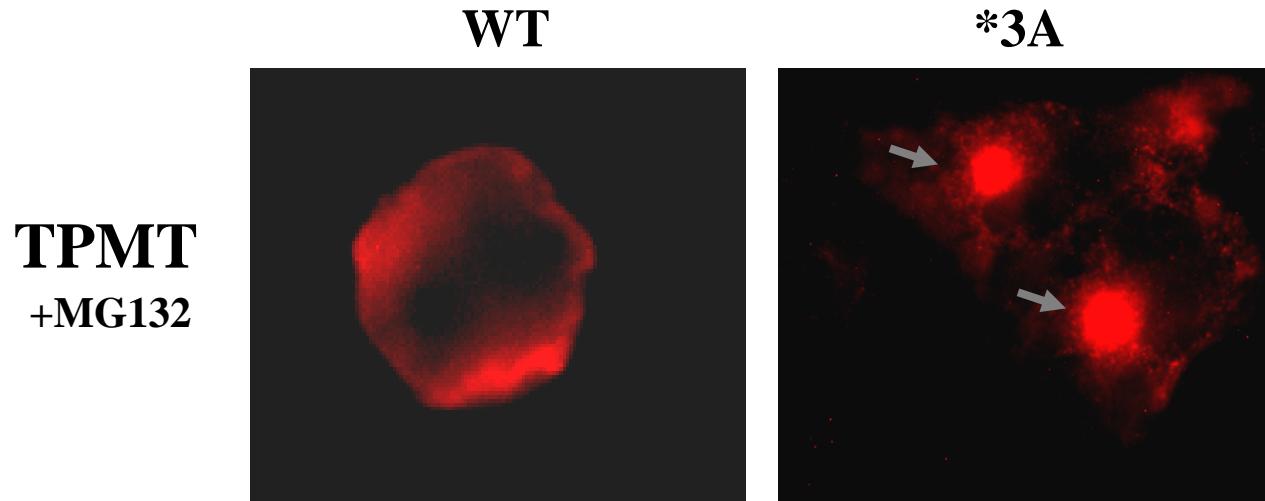
**G460A (Ala154 Thr) and A719G  
(Tyr240Cys) might result in  
TPMT misfolding and  
aggregation**

# Aggresome Formation



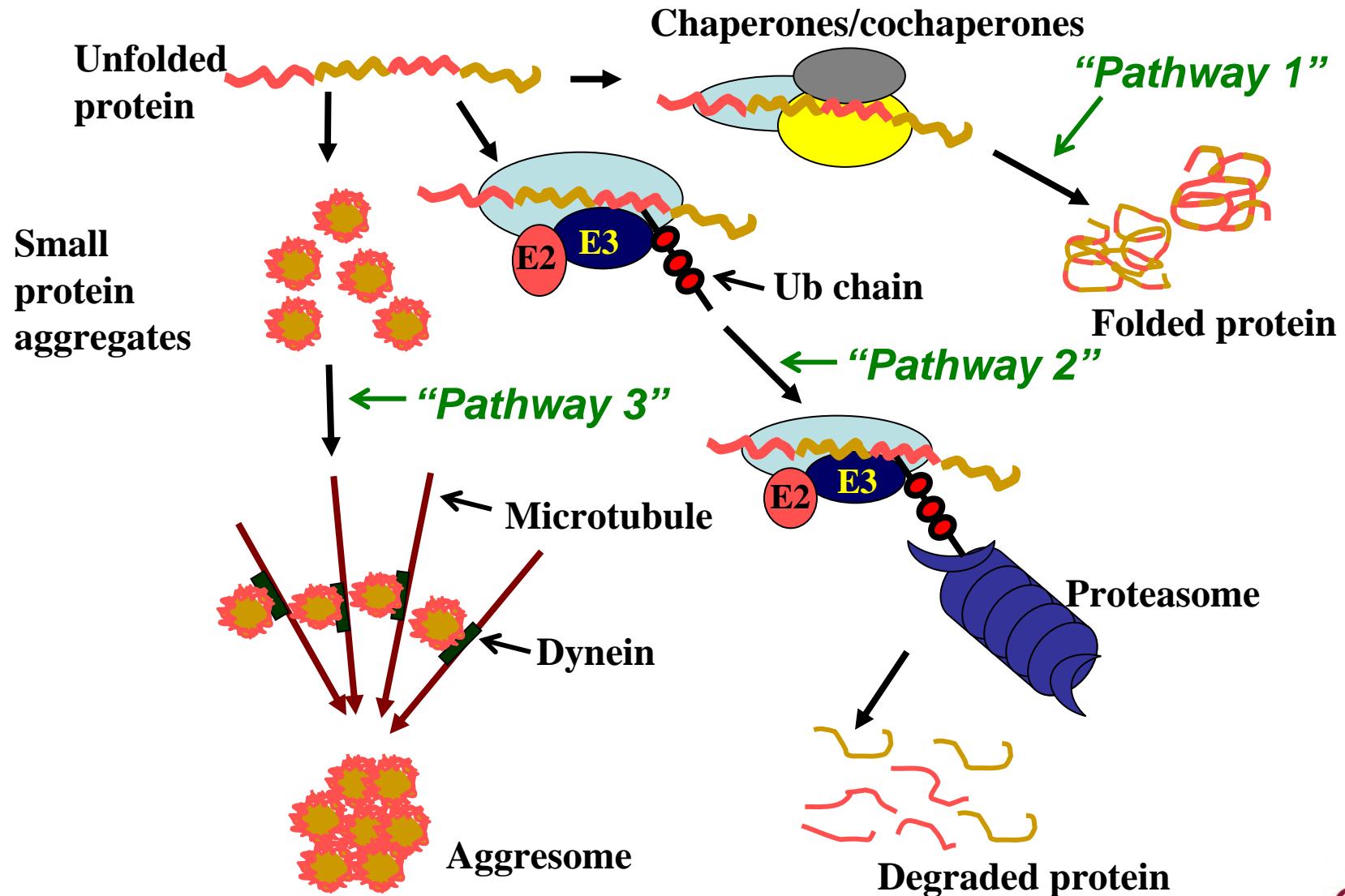
*Trends in Cell Biology*

# TPMT Aggresome Formation



Wang et al., PNAS 102:9394-99, 2005

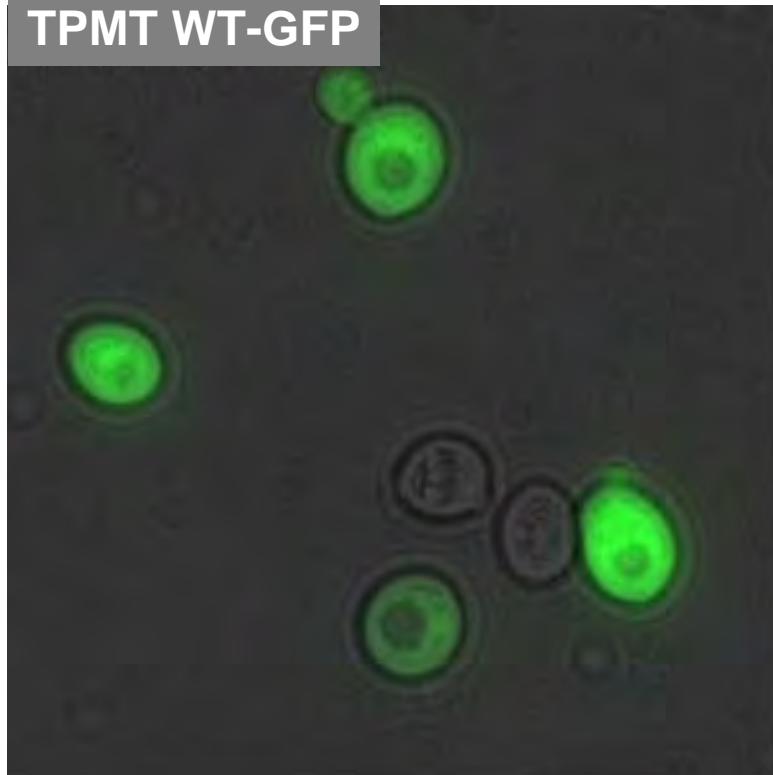
# Protein Folding, Degradation and Aggregation



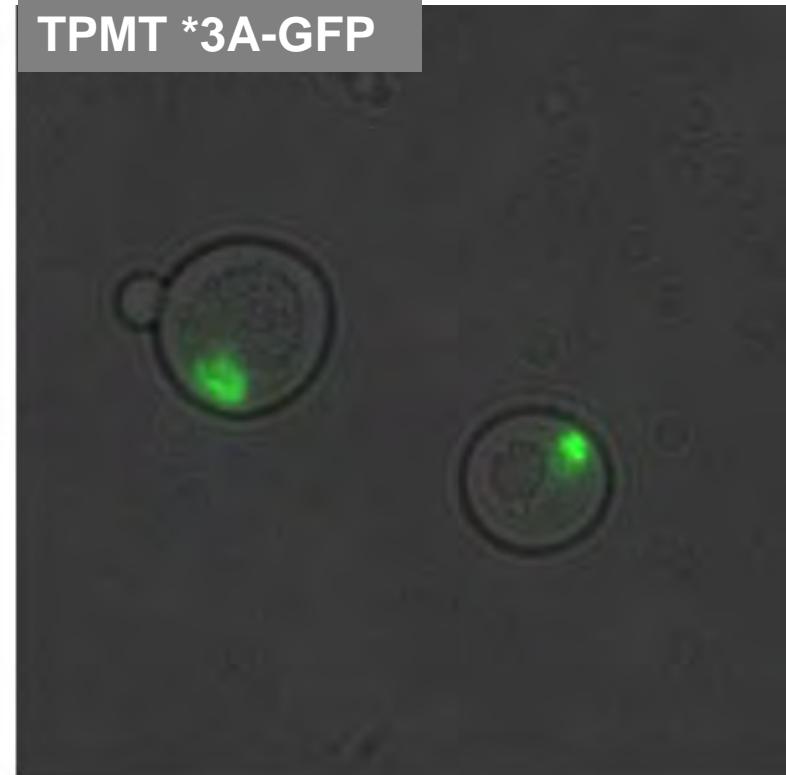
Wang and Weinshilboum, Oncogene 25:1629-38, 2006

# TPMT Allozymes in Yeast

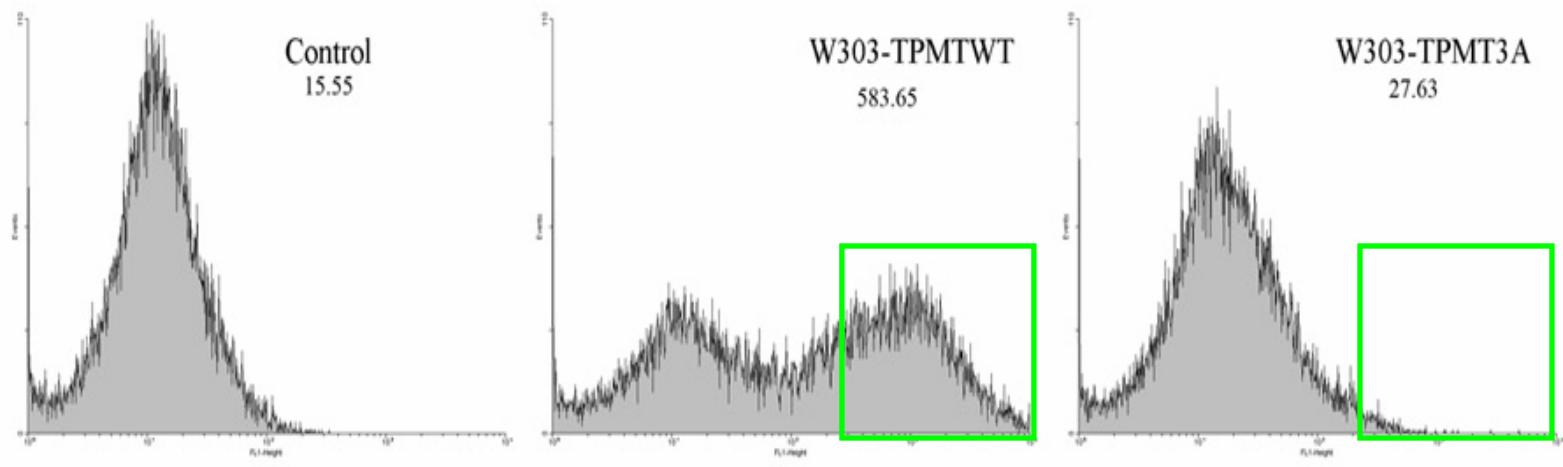
TPMT WT-GFP



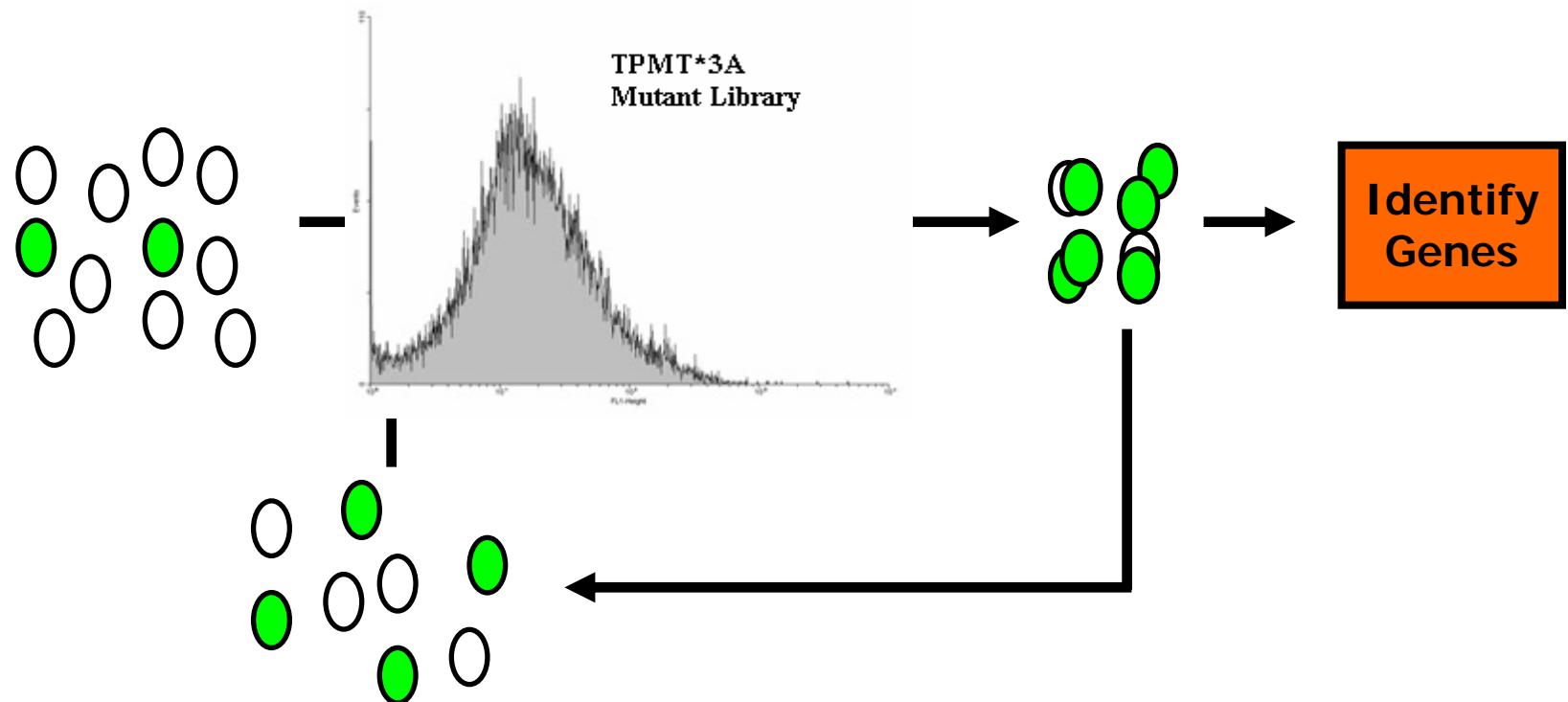
TPMT \*3A-GFP



# Yeast Flow Cytometry



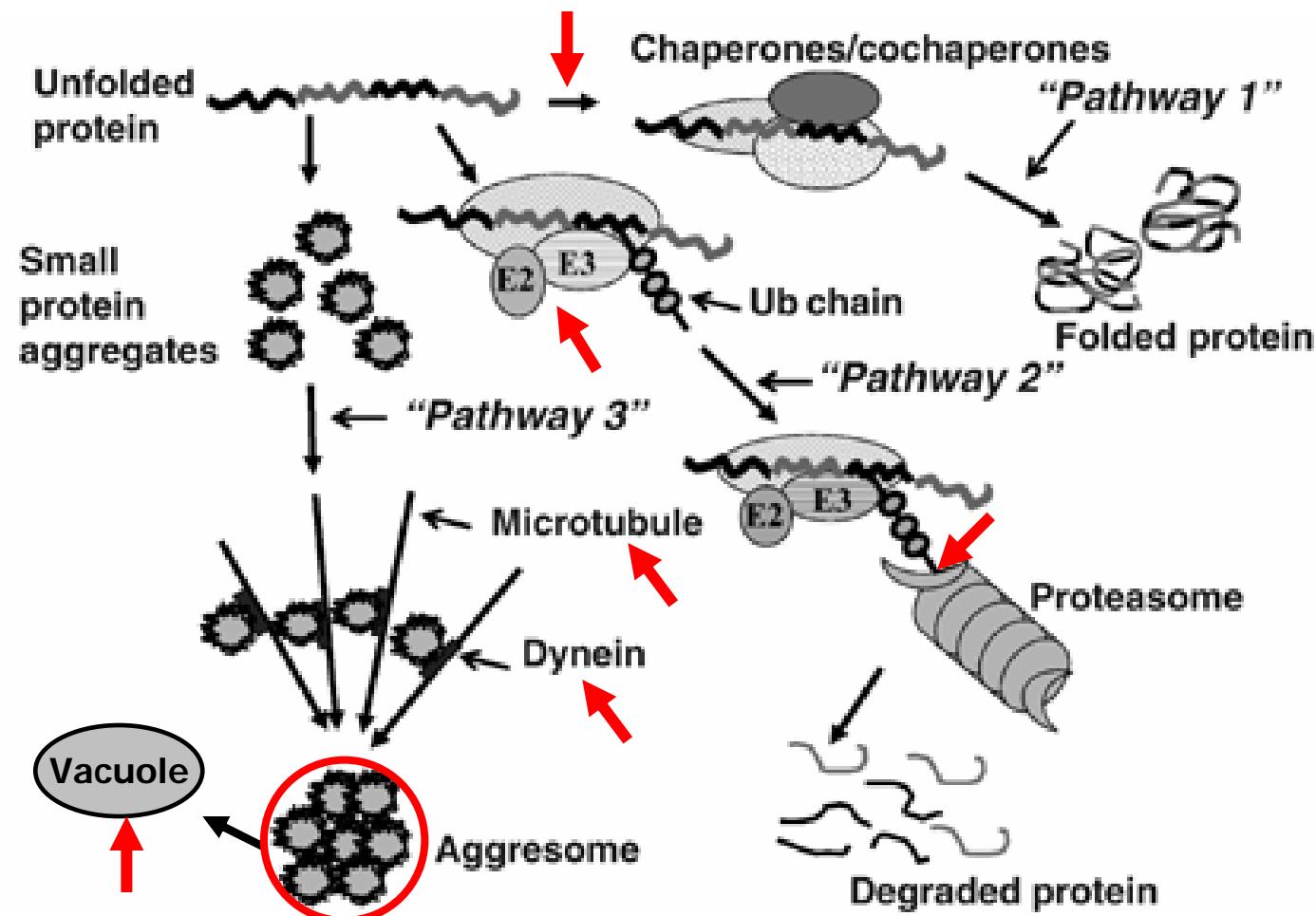
# Yeast TPMT Cell Sorting



# Sorting Results

- Thirty-one genes identified
- Ubiquitin mediated degradation pathway (E2, E3, proteasome)
- Chaperones (Hsp70)
- Microtubules (tubulin, dynein regulation, spindle pole body)
- Vesicle processes (trafficking, autophagy)

# Protein Folding, Degradation and Aggregation



# Pharmacogenomics

## Beyond Monogenic Pharmacogenomics

- Pathway-based
- Genome-wide association studies (GWA)
- Model systems – yeast, cell lines, etc.

# Pharmacogenomic Translational Studies

## Evolution

- One gene, one or a few SNPs
- One gene, intragene haplotypes
- PK and PD pathways and haplotypes
- Genome-wide association studies

# Pharmacogenomics

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- **Pharmacogenomic translation**
- Conclusions

# Pharmacogenetics-Pharmacogenomics

## FDA Hearings

## Pharmacogenetics and Drug Labeling

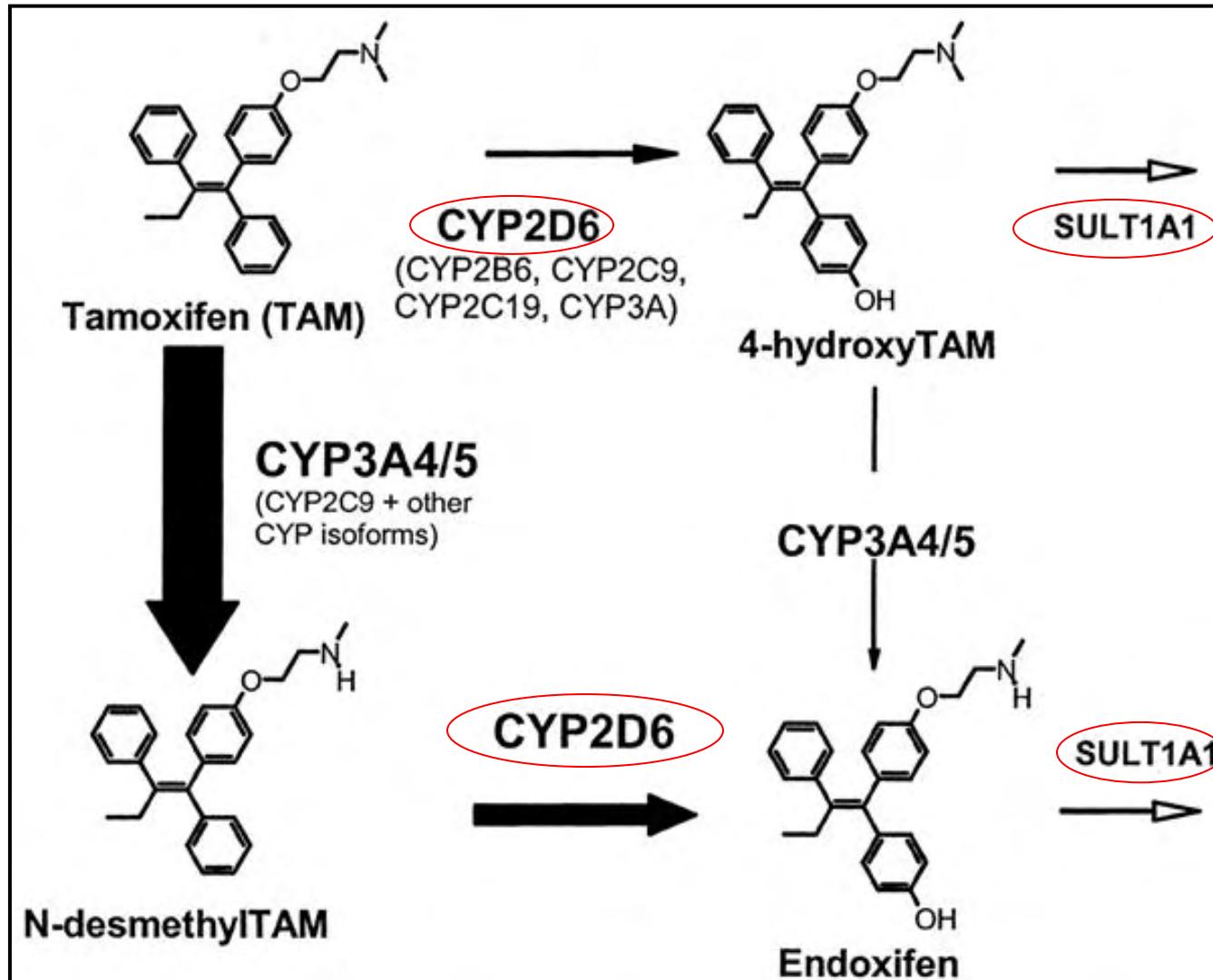
- Thiopurines – *TPMT*
- Irinotecan – *UGT1A1*
- Warfarin – *CYP2C9* and *VKORC1*
- Tamoxifen – *CYP2D6*

# Tamoxifen Pharmacogenetics

## Translational Pharmacogenomics



# Tamoxifen Biotransformation



Jin et al., J. Natl. Cancer Inst. 97:20-39, 2005.

# Post-Menopausal Breast Cancer Prospective Tamoxifen Trial *CYP2D6* Genotype and Outcomes

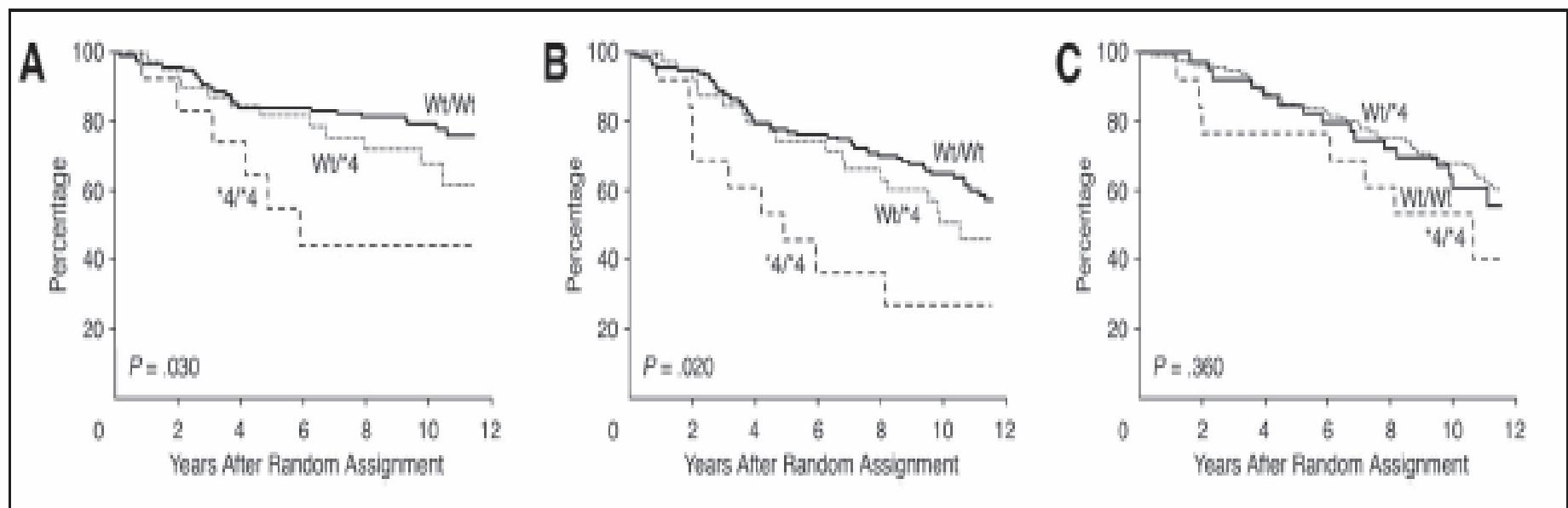


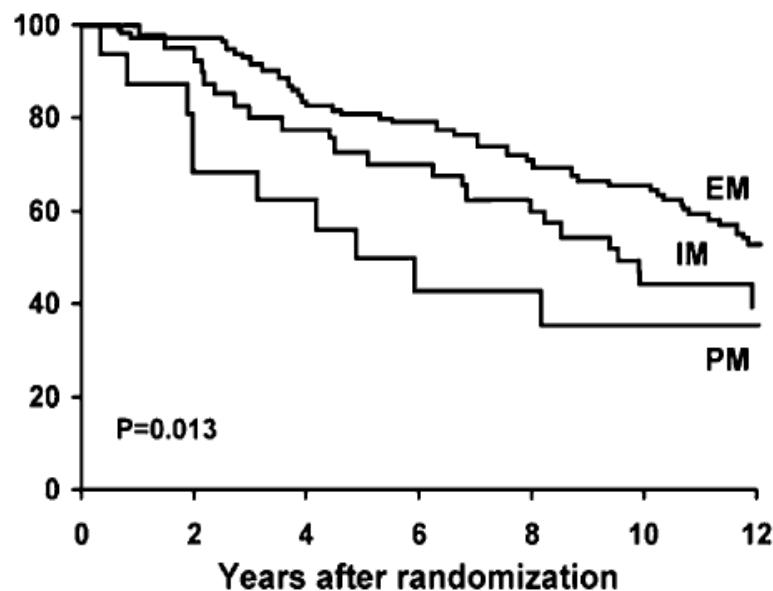
Fig 2. Kaplan-Meier estimates of (A) relapse-free time, (B) disease-free survival, and (C) overall survival for patients with the *CYP2D6\*4* genotype.

From Goetz M, Rae JM et al: *J Clin Oncol* 23:9312-9318, 2005.

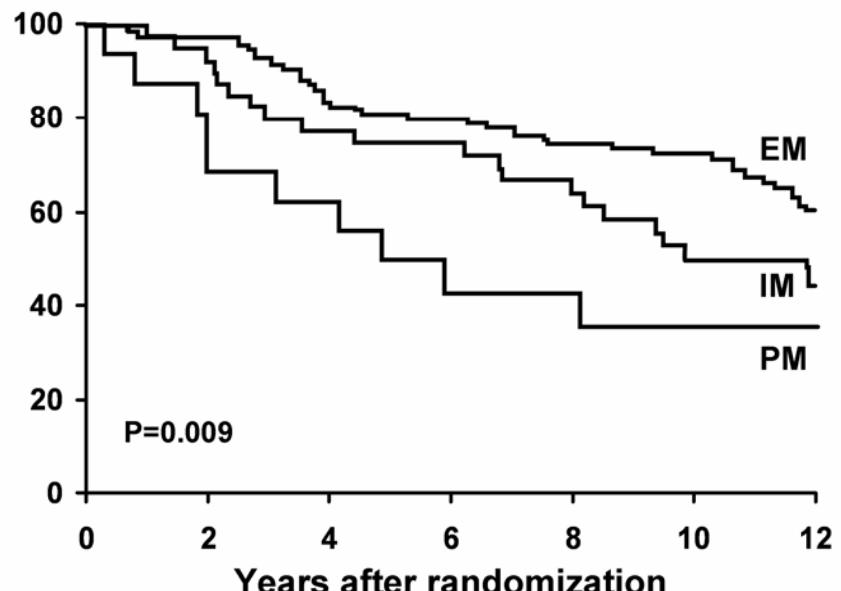
# Tamoxifen Pharmacogenomics

## Breast Cancer (190 Patients)

Relapse – Free Survival, %

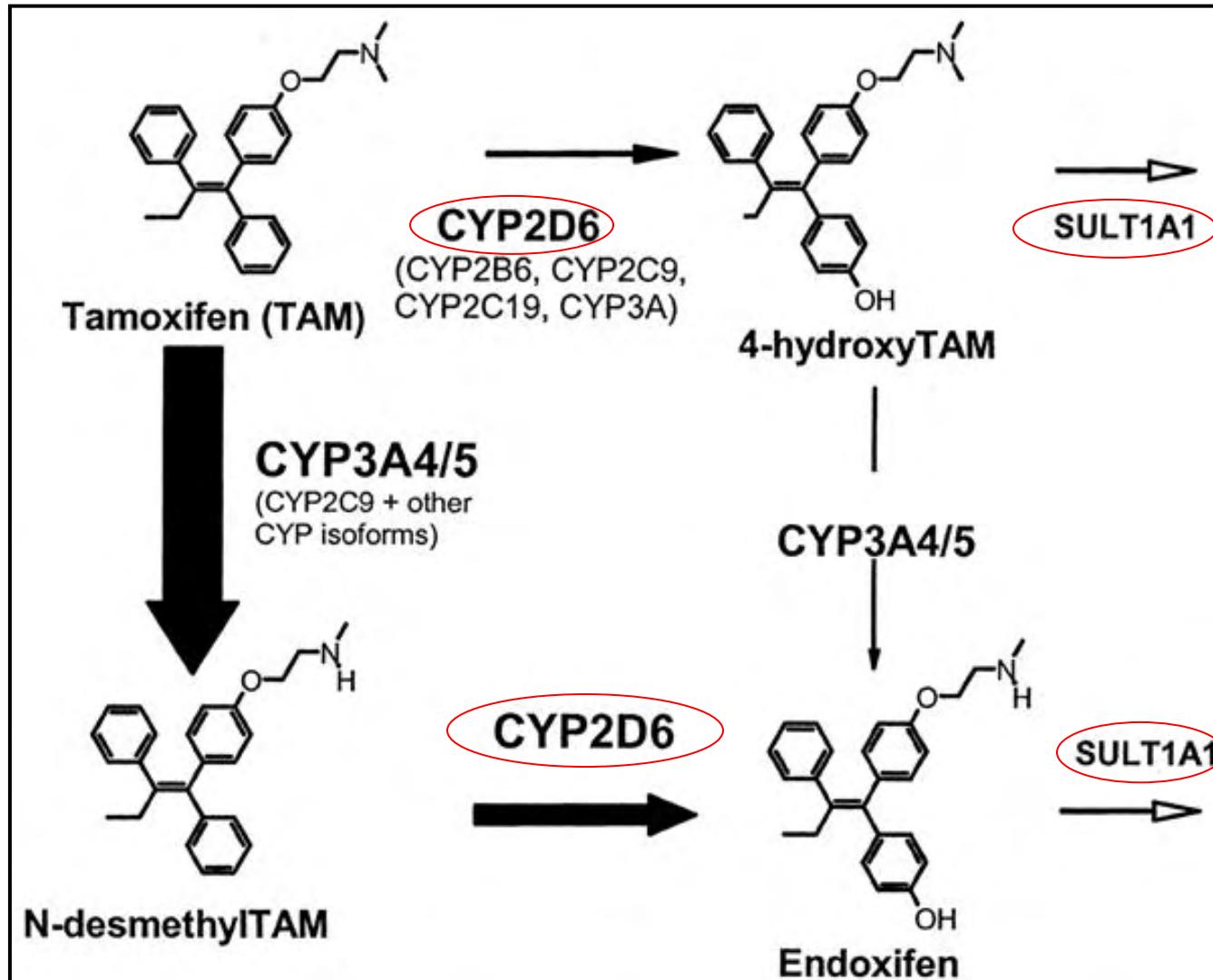


Disease – Free Survival



Goetz et al., Breast Cancer Res. Treat. 101:113-121, 2007.

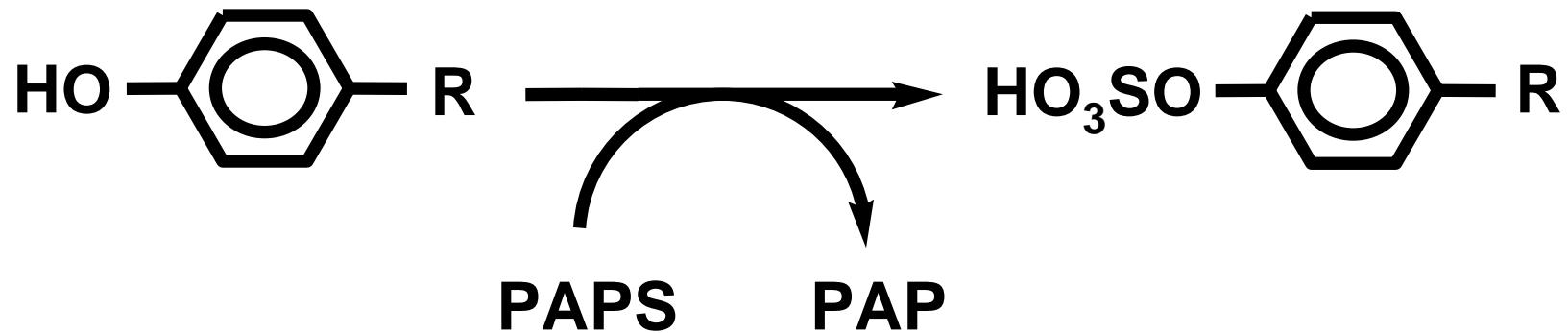
# Tamoxifen Biotransformation



Jin et al., J. Natl. Cancer Inst. 97:20-39, 2005.

# SULT Pharmacogenomics

## Phenol Sulfation

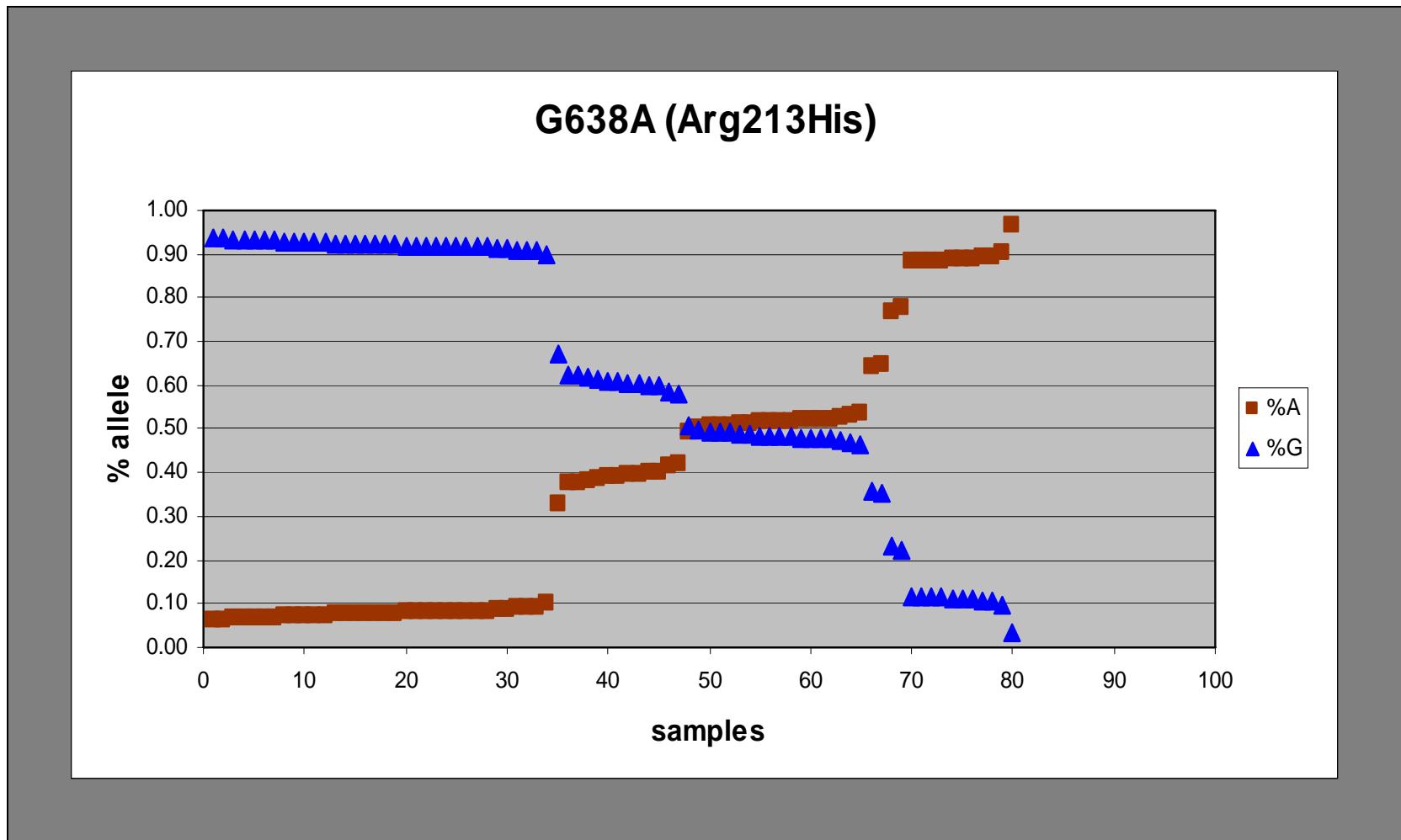


# Human SULT Isoforms

<u>SULT Gene</u>	<u>Chromosomal Location</u>	<u>Prototypic Substrate(s)</u>
<b>SULT1A1</b>		4-Nitrophenol and Estrogens
<b>SULT1A2</b>	16p11.2-12.1	Catecholamines
<b>SULT1A3</b>		
<b>SULT1A4</b>		
<b>SULT1C2</b>	2q11.2	4-Nitrophenol and Procarcinogens
<b>SULT1C4</b>		
<b>SULT1B1</b>	4q13	Thyroid hormones and 4-Nitrophenol
<b>SULT1E1</b>		Estrogens
<b>SULT2A1</b>		DHEA
<b>SULT2B1a</b>	19q13.3	Pregnenolone and DHEA
<b>SULT2B1b</b>		Cholesterol and DHEA
<b>SULT6B1</b>	2p22.3	Unknown
<b>SULT4A1</b>	22q13.1	Unknown

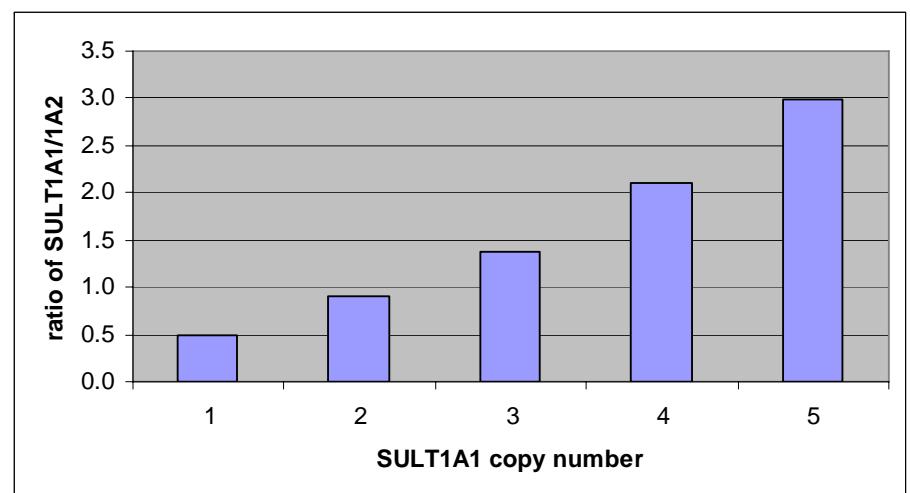
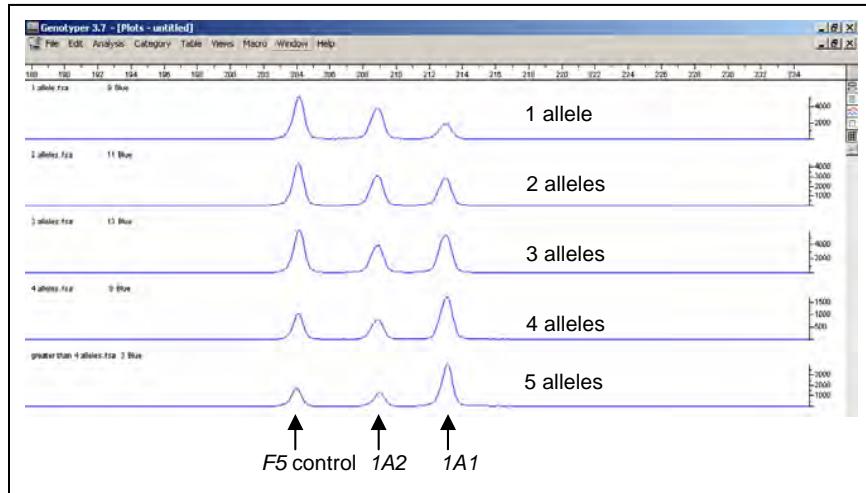
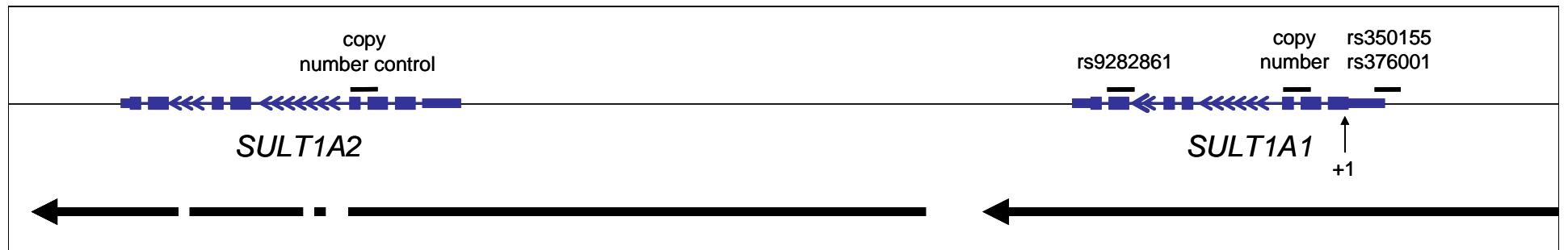
The phylogenetic tree on the left illustrates the evolutionary relationships between the various SULT genes. The tree branches into several main clades, with SULT1A1 and SULT1E1 being highlighted with red boxes around their respective gene names. Other genes are shown in blue.

# *SULT1A1* Pyrosequencing Results



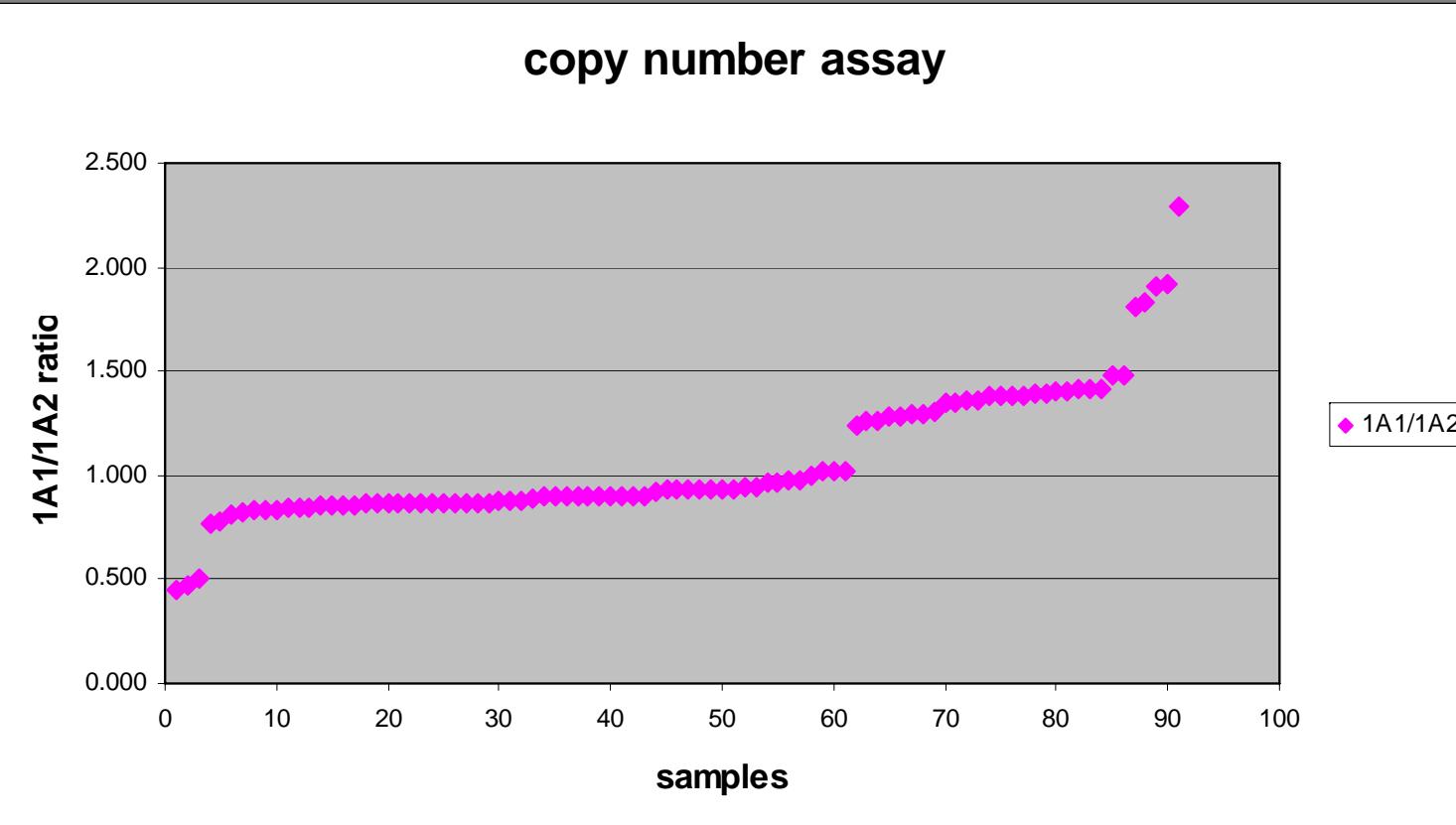
Hebbring et al., Human Mol. Genet. 16:463-470, 2006.

# *SULT1A1* Copy Number Assay



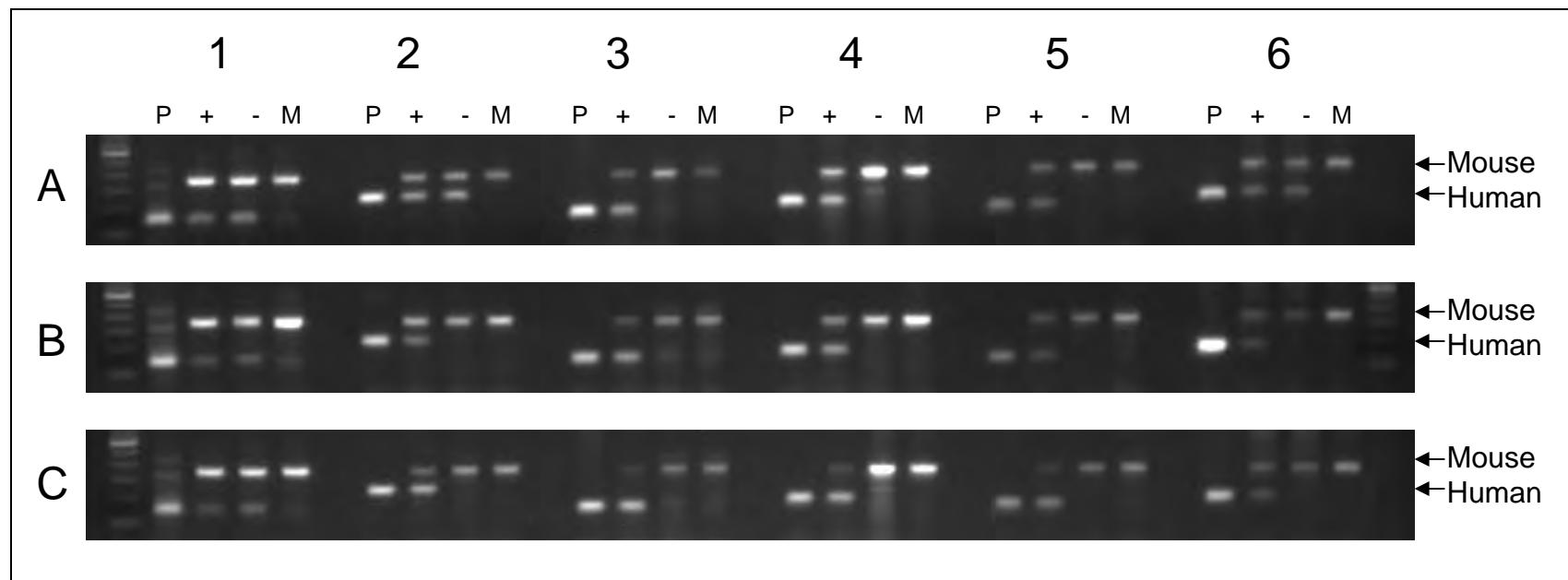
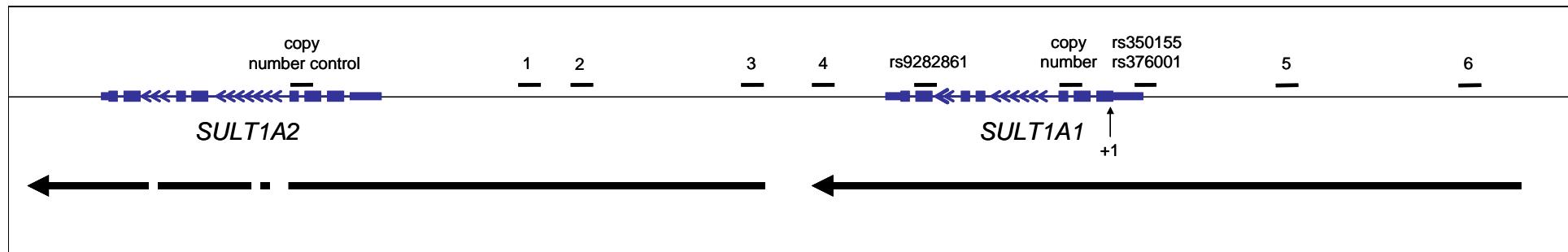
Hebbring et al., Human Mol. Genet. 16:463-470, 2006.

# *SULT1A1* Copy Number Assay



Hebbring et al., Human Mol. Genet. 16:463-470, 2006.

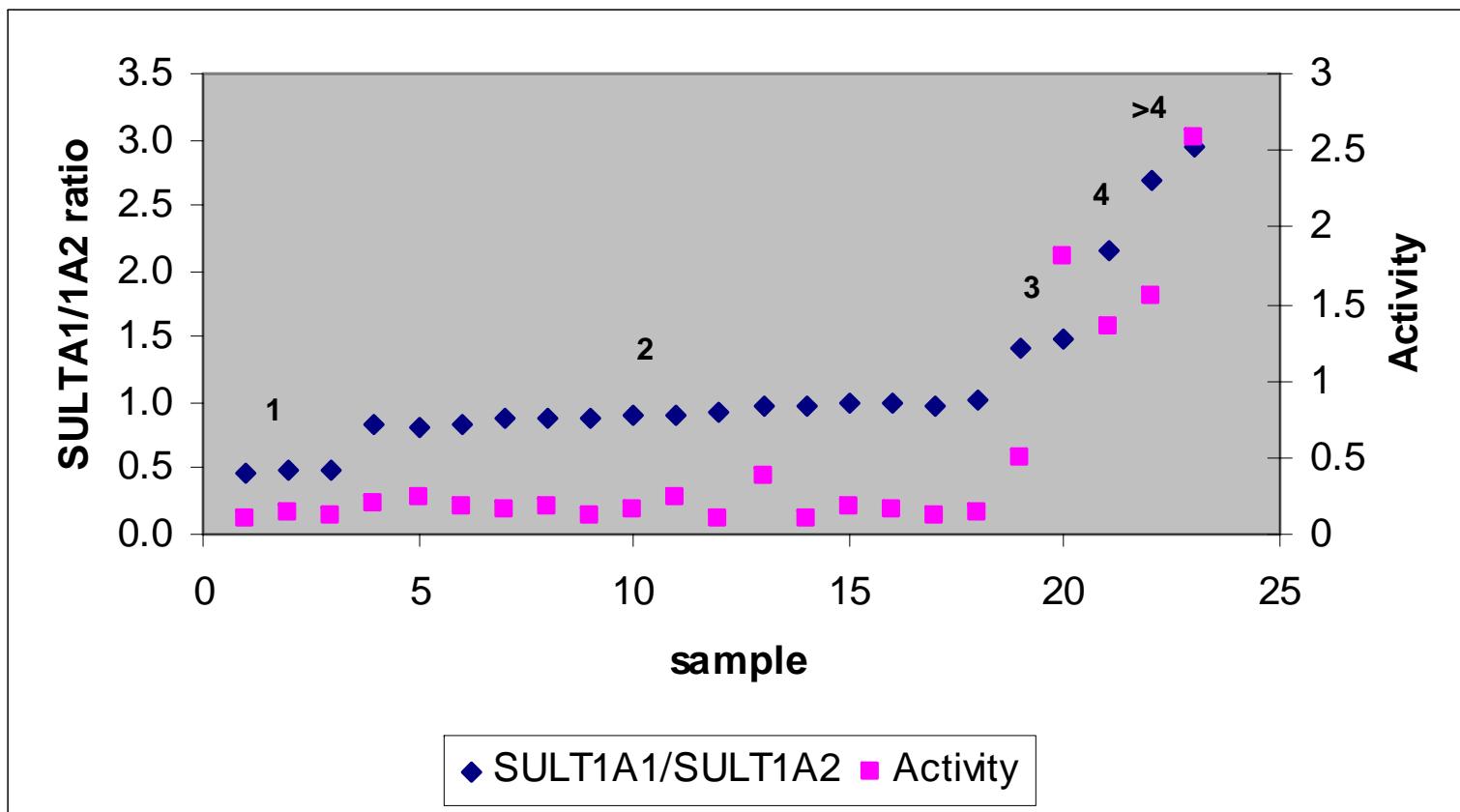
# *SULT1A1* Deletion Mapping



Hebbring et al., Human Mol. Genet. 16:463-470, 2006.

# SULT1A1 Activity vs Copy Number

## Platelet Samples n=23



p<0.0001

Hebbring et al., Human Mol. Genet. 16:463-470, 2006.

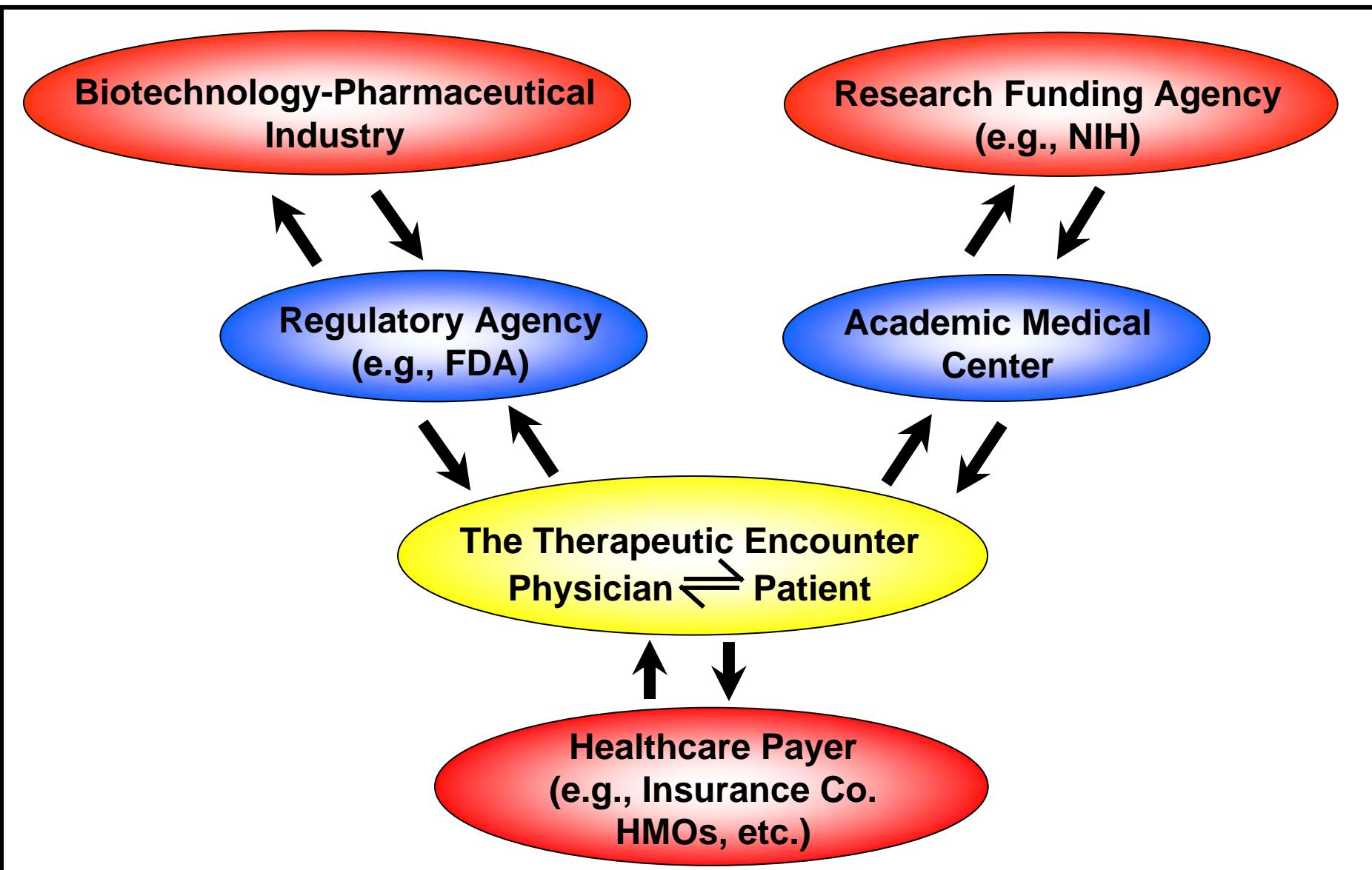
# Pharmacogenetics-Pharmacogenomics

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- **Irinotecan – *UGT1A1***
- **Warfarin – *CYP2C9* and *VKORC1***
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# Pharmacogenomic Clinical Translation

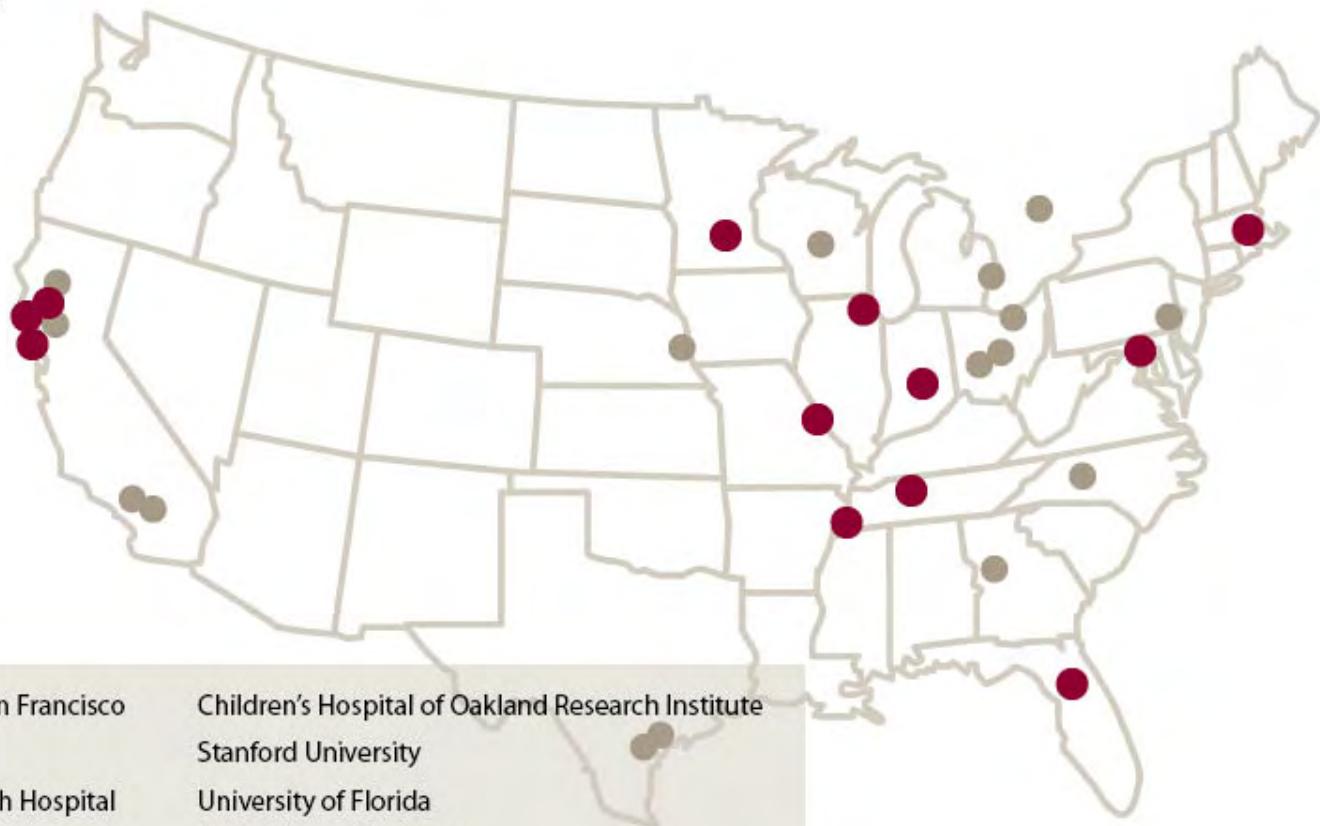


# Pharmacogenomics

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**NIGMS**  
**NHLBI**  
**NIDA**  
**NCI**  
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**NHGRI**  
**NLM**  
**ORWH**

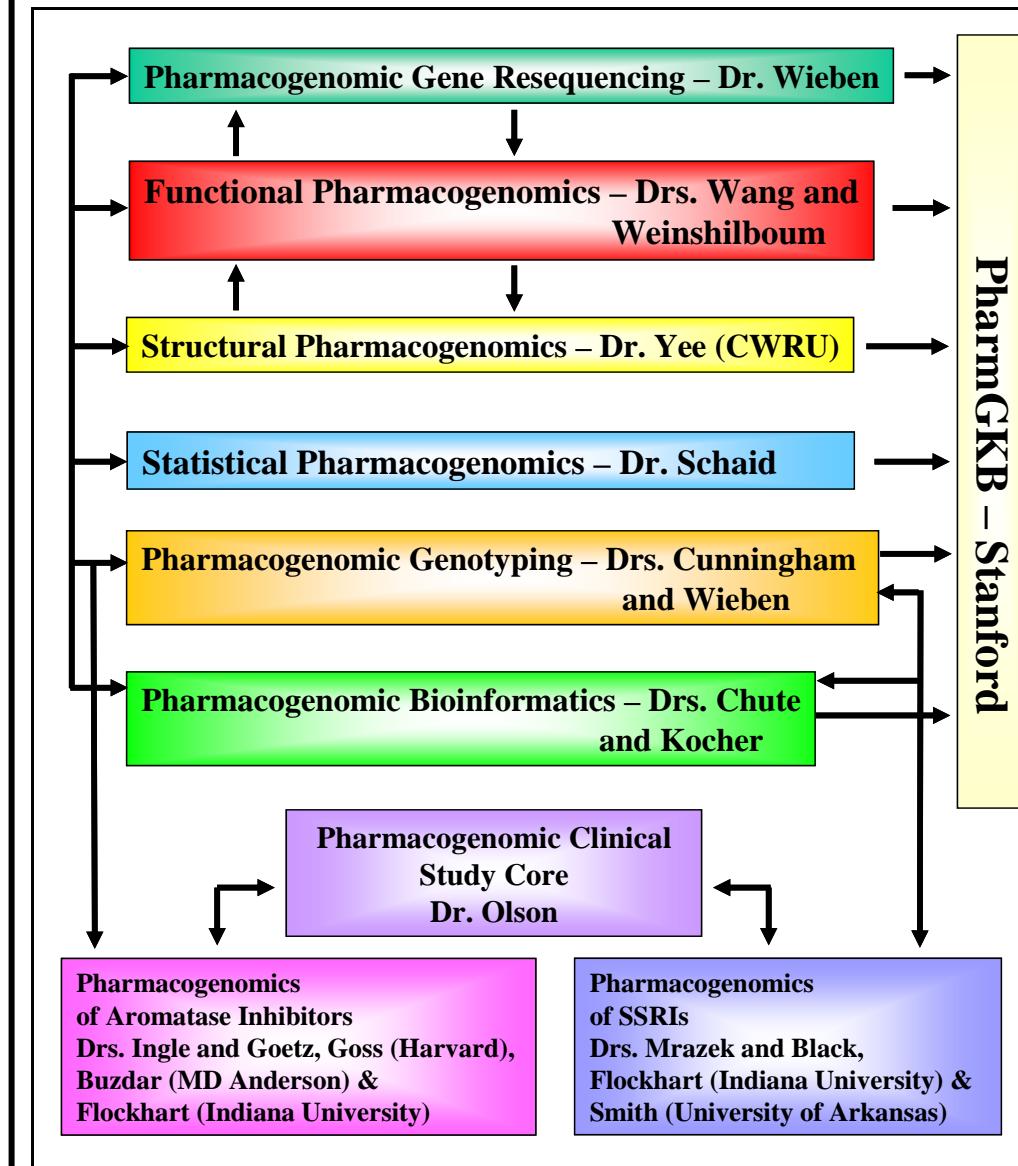


University of California, San Francisco  
University of Chicago  
St. Jude Children's Research Hospital  
Mayo Clinic  
Vanderbilt University  
Washington University

Children's Hospital of Oakland Research Institute  
Stanford University  
University of Florida  
University of Maryland  
Indiana University  
Brigham and Women's Hospital

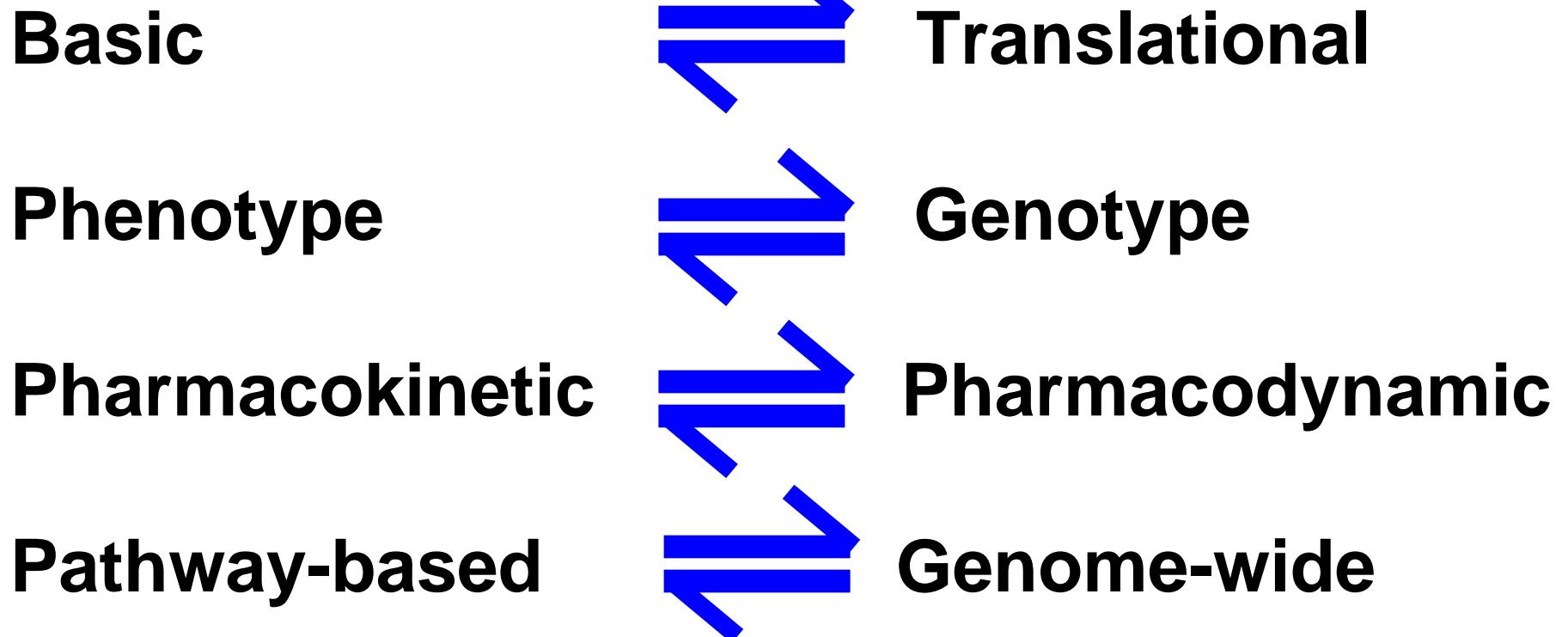
● Primary Investigator Site  
● Co-Investigator Site

# Mayo PGRN



# **Pharmacogenetics-Pharmacogenomics**

## **Complementary Approaches**



# **Pharmacogenetics-Pharmacogenomics**

**Discovery Science  
and  
Hypothesis-Based Science**



# Pharmacogenetics Coworkers

Araba Adjei

Ibrahim Aksoy

Pinar Aksoy

Saime Aksoy

Robert J. Anderson

Janel Baer

Blanka Boudikova-Girard

Norman Campbell

Edward Carlini

Monika Deininger

Joel Dunnette

Qiping Feng

Robert Freimuth

Todd Glauser

Chengtao Her

Sergio Hernandez

Michelle Hildebrandt

Ronald Honchel

Geoffrey Johnson

Julie Johnson

Yuan Ji

Krishna Kalari

Richard Keith

Mani Keshtgarpour

Adrian Kerremans

Annette Klumpp

Daniel Lee

Fang Li

Liang Li

Bonnie Maidak

Yvette Martin

Timothy Maus

Irene Moon

Ann Moyer

Gabriel Mwaluko

Diane Otterness

Patricia Pazmino

Randall Pearson

Linda Pelleymounter

Charles Preuss

Rebecca Raftogianis

Blanchard

Frederick Raymond

Christoph Reiter

Josephine Rini

Ezequiel Salavaggione

Alison Shield

Mary Scott

Susan Sladek

Rajah Sundarm

Carol Szumlanski

Bianca Thomae

Michael Thompson

Daniel Toft

Jon Van Loon

Randall Walker

Liewei Wang

Bill Watson

Thomas Wood

Lee Woodson

Luanne Wussow

Michael Xu

Lan Yan

Chen Yang

William Young Jr.

Jessica Zhang

Scott Hebring



# Mayo Pharmacogenomics Laboratory -- 2006

